1.   In a system including a legacy system producing clinical data for storage in a data repository, the clinical data having a format specific to the legacy system, a method for matching the clinical data to a standard of the clinical data before storing the clinical data in the data repository, the method comprising:

an act of receiving the clinical data from the legacy system at a health data dictionary;

an act of translating the clinical data by the health data dictionary such that the clinical data has a new format that is compatible with the standard;

an act of comparing the new format of the clinical data with the standard of the clinical data; and

when a match is found between the new format of the clinical data and the standard of the clinical data, an act of identifying one or more concept identifiers for the clinical data.

2.   A method as defined in claim 1, wherein the act of receiving the clinical data further comprises an act of receiving the clinical data through an interface engine, wherein the interface engine provides an interface code.

3.   A method as defined in claim 2, wherein the act of translating the clinical data further comprises an act of accessing the health data dictionary using the interface code.

4.   A method as defined in claim 1, wherein the act of translating the clinical data further comprises an act of identifying attributes of the clinical data.

5.   A method as defined in claim 4, wherein the act of identifying attributes further comprises an act of parsing the clinical data.

6.   A method as defined in claim 4, further comprising an act of identifying attributes from the clinical data, wherein the attributes correspond to attributes of the standard.

7.   A method as defined in claim 4, further comprising an act of using synonym tables to identify the attributes of the clinical data, wherein the synonym tables list equivalent expressions of the attributes.

8.   A method as defined in claim 4, further comprising an act of using relationship tables to define the clinical data.

9.   A method as defined in claim 1, further comprising an act of storing the standard format of the clinical data in the data repository, wherein the one or more concept identifiers are stored with the clinical data.

10.   A method as defined in claim 9, further comprising an act of retrieving the clinical data from the data repository.

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11.   A method as defined in claim 1, wherein the clinical data is laboratory results and wherein the standard format is Logical Observation Identifier Names and Codes.

12.   A computer program product having computer executable instructions for performing the acts recited in claim 1.

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13.   In a system including a legacy system providing clinical data including laboratory results to be stored in a data repository, wherein the laboratory results are in a format specific to the legacy system, a method for matching the clinical data including the laboratory results to a health data dictionary, the method comprising:

an act of loading standard laboratory results into the health data dictionary, wherein each standard laboratory result is associated with a unique concept identifier;

an act of creating standard relationship sets for each unique standard laboratory result, wherein the relationship sets establish relationships for attributes of each unique standard laboratory result;

an act of creating synonym tables for the attributes of the unique standard laboratory results;

an act of receiving the laboratory results at the health data dictionary;

an act of deriving attributes from the laboratory results using the synonym tables;

an act of generating a legacy relationship set for the laboratory results from the derived attributes; and

comparing the legacy relationship set with the standard relationship sets.

14.   A method as defined in claim 13, wherein the standard relationship sets identify attributes of each unique standard laboratory result.

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15.   A method as defined in claim 13, further comprising an act of determining if a new standard laboratory result should be added to the health data dictionary if an exact match is not found with the legacy laboratory result.

16.   A method as defined in claim 13, further comprising an act of comparing respective attributes of the legacy relationship table with the standard relationship tables.

17.   A method as defined in claim 13, further comprising an act of preventing matching inconsistencies using rules.

18.   A method as defined in claim 17, wherein the rules includes at least one of: frequency mapping; and suggesting a most likely match.

19.   A method as defined in claim 13, wherein the attributes include a component attribute, a property attribute, a time attribute, a system attribute, a scale attribute, and a method attribute.

20.   A method as defined in claim 13, further comprising an act of storing a matched laboratory result in the data repository, wherein the match laboratory result is normalized.

21.   A method as defined in claim 13, further comprising an act of manually matching laboratory results that do not match the standard laboratory results.

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22.   A computer program product having computer executable instructions for performing the acts recited in claim 12.

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23. In a system including a legacy transmitting legacy clinical information to a health data dictionary, a method for translating the clinical information to match a standard clinical information, the method comprising:

a step for creating standard sets of relationships for the standard clinical information in the health data dictionary;

a step for deriving legacy sets of relationships for the legacy clinical information; and

a step for comparing the legacy sets of relationships with the standard sets of relationships to identify an exact match for the legacy clinical information.

24. A method as defined in claim 23, wherein the step for creating standard sets of relationship further comprises a step for creating unique identifiers for each different code in the standard clinical information.

25. A method as defined in claim 24, wherein the step for creating standard sets of relationships further comprises:

a step for creating code relationship tables for each code, wherein the code relationship tables identify attributes of the standard clinical data; and

a step for creating attribute relationship tables for each code, wherein the attribute relationship tables identify independent values of the attributes of the standard clinical data.

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26. A method as defined in claim 25, wherein the step of deriving legacy sets of relationships further comprises a step for identifying independent values of the attributes using synonym tables, wherein the synonym tables contain synonyms for independent values.

27. A method as defined in claim 26, further comprising a step for entering the derived attributes in the legacy sets of relationships.

28. A method as defined in claim 23, further comprising a step for adding a new standard sets of relationships for legacy sets of relationships that do not match standard sets of relationships.

29. A method as defined in claim 23, further comprising a step for suggesting a match when the legacy sets of relationships partially match the standard sets of relationships.

30. A method as defined in claim 23, wherein the standard sets of relationships comply with Logical Observation Identifier Names and Codes.

31. A computer program product having computer executable instructions for performing the steps recited in claim 23.

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